RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09 856,812B

Source: 1F W16

Date Processed by STIC: 11/07/2005

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       RAW SEQUENCE LISTING              DATE: 11/07/2005
       PATENT APPLICATION:  US/09/856,812B      TIME: 12:22:40

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  3  <110> APPLICANT: Huang, Lan-Qing
  4         Van Pel, Aline
  5         Brasseur, Francis
  6         De Plaen, Etienne
  7         Boon, Thierry
  9  <120> TITLE OF INVENTION: Tumour Rejection Antigens
 11  <130> FILE REFERENCE: L0461.70115US00
 13  <140> CURRENT APPLICATION NUMBER: US 09/856,812B
-> 14 <141> CURRENT FILING DATE: 2001-09-07
 16  <150> PRIOR APPLICATION NUMBER: GB 9826143.1
 17  <151> PRIOR FILING DATE: 1998-11-27
 19  <160> NUMBER OF SEQ ID NOS: 57
 21  <170> SOFTWARE: PatentIn Ver. 3.2
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 33               20                  25                  30
 35  Ala Val Glu Glu Asp Ala Ser Ser Ser Thr Ser Thr Ser Ser Ser Phe
 36           35                  40                  45
 38  Pro Ser Ser Phe Pro Ser Ser Ser Ser Ser Ser Ser Ser Cys Tyr
 39       50                  55                  60
 41  Pro Leu Ile Pro Ser Thr Pro Glu Glu Val Ser Ala Asp Asp Glu Thr
 42   65                  70                  75                  80
 44  Pro Asn Pro Pro Gln Ser Ala Gln Ile Ala Cys Ser Ser Pro Ser Val
 45               85                  90                  95
 47  Val Ala Ser Leu Pro Leu Asp Gln Ser Asp Glu Gly Ser Ser Ser Gln
 48               100                 105                 110
 50  Lys Glu Glu Ser Pro Ser Thr Leu Gln Val Leu Pro Asp Ser Glu Ser
 51           115                 120                 125
 53  Leu Pro Arg Ser Glu Ile Asp Glu Lys Val Thr Asp Leu Val Gln Phe
 54       130                 135                 140
 56  Leu Leu Phe Lys Tyr Gln Met Lys Glu Pro Ile Thr Lys Ala Glu Ile
 57  145                 150                 155                 160
 59  Leu Glu Ser Val Ile Lys Asn Tyr Glu Asp His Phe Pro Leu Leu Phe
 60               165                 170                 175
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 63               180                 185                 190
 65  Glu Val Asp Pro Thr Gly His Ser Phe Val Leu Val Thr Ser Leu Gly
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66          195                 200                 205
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72 225                 230                 235                 240
74 Thr Pro Glu Glu Val Ile Trp Glu Ala Leu Asn Met Met Gly Leu Tyr
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77 Asp Gly Met Glu His Leu Ile Tyr Gly Glu Pro Arg Lys Leu Leu Thr
78                 260                 265                 270
80 Gln Asp Trp Val Gln Glu Asn Tyr Leu Glu Tyr Arg Gln Val Pro Gly
81       275                 280                 285
83 Ser Asp Pro Ala Arg Tyr Glu Phe Leu Trp Gly Pro Arg Ala His Ala
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86 Glu Ile Arg Lys Met Ser Leu Leu Lys Phe Leu Ala Lys Val Asn Gly
87 305                 310                 315                 320
89 Ser Asp Pro Arg Ser Phe Pro Leu Trp Tyr Glu Glu Ala Leu Lys Asp
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92 Glu Glu Glu Arg Ala Gln Asp Arg Ile Ala Thr Thr Asp Asp Thr Thr
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117 Thr Leu Glu Glu Val Thr Asp Ser Gly Ser Pro Ser Pro Pro Gln Ser
118       50                 55                 60
120 Pro Glu Gly Ala Ser Ser Ser Leu Thr Val Thr Asp Ser Thr Leu Trp
121 65                 70                 75                 80
123 Ser Gln Ser Asp Glu Gly Ser Ser Ser Asn Glu Glu Glu Gly Pro Ser
124                 85                 90                 95
126 Thr Ser Pro Asp Pro Ala His Leu Glu Ser Leu Phe Arg Glu Ala Leu
127                 100                 105                 110
129 Asp Glu Lys Val Ala Glu Leu Val Arg Phe Leu Leu Arg Lys Tyr Gln
130           115                 120                 125
132 Ile Lys Glu Pro Val Thr Lys Ala Glu Met Leu Glu Ser Val Ile Lys
133       130                 135                 140
135 Asn Tyr Lys Asn His Phe Pro Asp Ile Phe Ser Lys Ala Ser Glu Cys
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138 Met Gln Val Ile Phe Gly Ile Asp Val Lys Glu Val Asp Pro Ala Gly
139                 165                 170                 175
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141 His Ser Tyr Ile Leu Val Thr Cys Leu Gly Leu Ser Tyr Asp Gly Leu
142             180             185             190
144 Leu Gly Asp Asp Gln Ser Thr Pro Lys Thr Gly Leu Leu Ile Ile Val
145         195             200             205
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182 tggtttaagg aagtggcctc agatcaacaa agggagggtc ccaggtcgta tcaggcatca 1140
183 agaagaggac caagcaggct cctcacccca gtacacatgg acccagctga atatggccac 1200
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195 ggccacttac accctccctc tctccccagg cctgtgggtc cccatcgccc aagtcctgcc 1920
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206 Gly Leu Glu Gly Ala Gln Ala Pro Leu Ala Val Glu Glu Asp Ala Ser
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209 tca tcc act tcc acc agc tcc tct ttt cca tcc tct ttt ccc tcc tcc   2119
210 Ser Ser Thr Ser Thr Ser Ser Ser Phe Pro Ser Ser Phe Pro Ser Ser
211 40                  45                  50                  55
213 tcc tct tcc tcc tcc tcc tcc tgc tat cct cta ata cca agc acc cca   2167
214 Ser Ser Ser Ser Ser Ser Ser Cys Tyr Pro Leu Ile Pro Ser Thr Pro
215                 60                  65                  70
217 gag gag gtt tct gct gat gat gag aca cca aat cct ccc cag agt gct   2215
218 Glu Glu Val Ser Ala Asp Asp Glu Thr Pro Asn Pro Pro Gln Ser Ala
219                 75                  80                  85
221 cag ata gcc tgc tcc tcc ccc tcg gtc gtt gct tcc ctt cca tta gat   2263
222 Gln Ile Ala Cys Ser Ser Pro Ser Val Val Ala Ser Leu Pro Leu Asp
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225 caa tct gat gag ggc tcc agc agc caa aag gag gag agt cca agc acc   2311
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227     105                 110                 115
229 cta cag gtc ctg cca gac agt gag tct tta ccc aga agt gag ata gat   2359
230 Leu Gln Val Leu Pro Asp Ser Glu Ser Leu Pro Arg Ser Glu Ile Asp
231 120                 125                 130                 135
233 gaa aag gtg act gat ttg gtg cag ttt ctg ctc ttc aag tat caa atg   2407
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238 Lys Glu Pro Ile Thr Lys Ala Glu Ile Leu Glu Ser Val Ile Lys Asn
239                 155                 160                 165
241 tat gaa gac cac ttc cct ttg ttg ttt agt gaa gcc tcc gag tgc atg   2503
242 Tyr Glu Asp His Phe Pro Leu Leu Phe Ser Glu Ala Ser Glu Cys Met
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246 Leu Leu Val Phe Gly Ile Asp Val Lys Glu Val Asp Pro Thr Gly His
247     185                 190                 195
249 tcc ttt gtc ctt gtc acc tcc ctg ggc ctc acc tat gat ggg atg ctg   2599
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253 agt gat gtc cag agc atg ccc aag act ggc att ctc ata ctt atc cta   2647
254 Ser Asp Val Gln Ser Met Pro Lys Thr Gly Ile Leu Ile Leu Ile Leu
255                 220                 225                 230
257 agc ata atc ttc ata gag ggc tac tgc acc cct gag gag gtc atc tgg   2695
258 Ser Ile Ile Phe Ile Glu Gly Tyr Cys Thr Pro Glu Glu Val Ile Trp
259                 235                 240                 245
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267       265                    270                    275
269 tac ctg gag tac cgg cag gtg cct ggc agt gat cct gca cgg tat gag  2839
270 Tyr Leu Glu Tyr Arg Gln Val Pro Gly Ser Asp Pro Ala Arg Tyr Glu
271 280                    285                    290                    295
273 ttt ctg tgg ggt cca agg gct cat gct gaa att agg aag atg agt ctc  2887
274 Phe Leu Trp Gly Pro Arg Ala His Ala Glu Ile Arg Lys Met Ser Leu
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277 ctg aaa ttt ttg gcc aag gta aat ggg agt gat cca aga tcc ttc cca  2935
278 Leu Lys Phe Leu Ala Lys Val Asn Gly Ser Asp Pro Arg Ser Phe Pro
279                315                    320                    325
281 ctg tgg tat gag gag gct ttg aaa gat gag gaa gag aga gcc cag gac  2983
282 Leu Trp Tyr Glu Glu Ala Leu Lys Asp Glu Glu Glu Arg Ala Gln Asp
283                330                    335                    340
285 aga att gcc acc aca gat gat act act gcc atg gcc agt gca agt tct  3031
286 Arg Ile Ala Thr Thr Asp Asp Thr Thr Ala Met Ala Ser Ala Ser Ser
287       345                    350                    355
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-> 291 360                    365                    370
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VERIFICATION SUMMARY   DATE: 11/07/2005
PATENT APPLICATION: **US/09/856,812B**   TIME: 12:22:41

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163 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:5